Figure S1
AluYconsensus
Human-BLAT
Chimp-BLAT
Pan paniscus(I)
Pan paniscus(C)
Gorilla gorilla
Orangutan-BLAT
Susie Ref.
GM06213A
GM04272A
NG12256
NG06209
AG05252A
KB4204
KB5404
KB5405
T G G T T A T T T G T T T T G G G G A A G TAACATTTTGCTTAATAGTGGTTCAAAAT 0
50
T G G T T A T T T G T T T G G T T A T T T G T T
TG G T T A T T T GT G TTATTTGTTCTGGGGAAGTAACATTTTGCTTAATAGTGGTTAAAAAT G G T TA T T T G T T C T G G G G A A G T A A CATTTTGCTCAATAGTGGTTCAAAATA50 T G G T T A T T T G T T T T G G G G A A G TAACATTTTGCTTAATAGTGGTTCAAAATM50 T G G T TA T T T G T T T T G G G G A A G TAACATTTTGCTTAATAGTGGTTCAAAAT 50 TGGTTATTTGTTTTGGGGAAGTAACATTTTGCTTAATAGTGGTTCAAAATM50 T G G T T A T T T G T T C T G G G G A A G TAACATTTTGCTTAATAGTGGTTCAAAATM50 T G G T TA T T T G T T T T G G G GAA G TAACATTTTGCTTAATAGTGGTTCAAAATM50 TGGTTATTTGTTTTGGGGAAGTAACATTTTGCTTAATAGTGGTTCAAAATA50 TGGTTATTTGTTTTGGGGAAGTAACATTTTGCTTAATAGTGGTTCAAAATM50 TGGTTATTTGTTTTGGGGAAGTAACATTTTGCTTAATAGTGGTTCAAAATM50 T G G T TATTTGTTTTGGGGAAGTAACATTTTGCTTAATAGTGGTTCAAAATM50
TGGTTATTTGTTTTGGGGAAGTAACATTTTGCTTAATAGTGGTTCAAAATA50

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#### Abstract

TGGTGTTTCGTTGTCATCAAATCAGTTACACATACACAAGCTG TGGTGTTC GTTGTCATCAAATCAGTTACACATACACAAGCTG GGTGTTCGTTGTCATCAAATCAGTTACACATACACAAGCTG T G G T G T T C G T T G T C A T CAAATCAGTTACACATACACAAGCTG TGGTGTTCGTTGTCATCAAATCAGTTACACATACACAAGCTG T G G T G T T C C T T G T CATCAAATCAGTTACATATACACAAGCTG T G G T G T T C C T T G T C A T C A A A T C A G T T A C A TATACACAA G T G T G G T G T TCCTTGTCATCAAATCAGTTACATATACACAACCTG TGGTGTTCCTTGTCATCAAATCAGTTACATATACACAAGCTG T G G T G T T C C T T G T C A T C A A A T C A G T T A C A TATACACAA G C T G T G G T G T T C C T T G T C A T C A A A T C A G T T A C A T A T A C A C A A G C T G TGGTGTTCCTTGTCATCAAATCAGTTACATATACACAAGCTG TGGTGTTCCTTGTCATCAAATCAGTTACATATACACAAGCTG T G G T G T T C C T T G T C A T C A A A TCAGTTACATATACACAAGCTG T G G T G T TCCTTGTCATCAAATCAGTTACATATACACAACCTG


 100 100 100 100
100 100
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102030 40

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KB4204
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G GCCGGGCGCGGTGGCTCACGCCTGTAATCCCAGCACTTT 40

| - | - | - | - | - | - | - | - | - | - | $G$ |
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| $\mathbf{C}$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $\mathbf{C}$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $\mathbf{C}$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $C$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $C$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $T$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |
| $C$ | $A$ | $G$ | $G$ | $T$ | $C$ | $T$ | $G$ | $T$ | $T$ | $G$ |


 $C \subset G G G T G C G G T G G C T C A C G C G T G T A A T C C A A C C A C T T T 150$ $C \subset G G G T G C G G T G G C T C A C G C G T G T A A T C C C A G C A C T T T 150$ A C C G G G C G C G G T G G C T C A C G C C T G T A A T C C C A G C A C T T C 150 $G G C C G G G C G C G G T G G C T C A C G C C T G T A A T C C C A G C A C T T T 150$ G GCCGGGCGCGGTGGCTCACGCCTGTAATCCCAGCACTTT150 G GCC G G G C G C G G T G G C T C A C G C C T G TAA T C C C A G C A C T T T 150

 G GCCGGGCGCGGTGGCTCACGCCTGTAATCCCAGCACTTT150
 G GCC C G G C G C G G T G G C T C A C G C C T G TAA T C C C A G C A C T T T 150 G GCCGGGCGCGGTGGCTCACGCCTGTAATCCAACCACTTT150 $G G C \subset G G G C G C G G T G G C T C A C G C C T G T A A T C C C A G C A C T T T 150$

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Gorilla gorilla
Orangutan-BLAT
Susie Ref.
GM06213A
GM04272A
NG12256
NG06209
AG05252A
KB4204
KB5404
KB5405

GGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG 90 G G GAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 G G GCGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 GGGCGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 GGGCGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 C G GAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 GGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 GGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 GGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 G G G A G GCC G A G G C G G G C A G A T C A C G A G G T CA G GA GATC G A G A C C A T C C T G 200 G G GAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 G G GAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 GGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 G G G A G G C C G A G G C G G G C G G A T C A C G A G G T C A G G A G A T C G A G A C C A T C C T G 200 G G G A G GCC G A G GCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200 GGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTG200
$\qquad$ 120

130
140
G C TAACACG GTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCGG140 GCTTACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCTG250 GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCTG250 GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCTG250 GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCTG250 GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCTG250

GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCTG250 GCTAACACGGCGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCC GCTAACACGGTGAAACCCCGTCTCTACTAAAAAATACAAAAAATTAGCC GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCC GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCC GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCC GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCC GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCC G C TAACACG GT GAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCC GCTAACACGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCC $150 \quad 160$ $\begin{array}{llllll}150 & 160 & 170 & 180 & 190\end{array}$ $\begin{array}{llllll}150 & 160 & 170 & 180 & 190\end{array}$

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G 250

GC G T G G T G GCGGGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGG190 GCAA G G T G GCGGGCGCCTGTAGTCCCAGCTACGCGGGAGGCTGAGGCAGGY00

## G C A

G C A
G C A G C A G C G G C G
G C G
G C G
G C G
G C G A
G C G
G C G
G C G
G C G G G T G G C G G G C G C C T G T A G TC C CA G C TC C TC G G G A G G C T G A G G C A G G 300 G GTGGCGGGCGCCTGTAGTCCCAGCTCCTCGGGAGGCTGAGGCAGA 300 G G T G GC G G GCGCCTGTAGTCCCAGCTCCTCGGGAGGCTGAGGCAGA300 GGTGGC G G GCGCCCTGTAGTCCCAGCTCCTC G G GAAGGCTGAGGCAGG300 GGTGGC G G GTGGCCTGTAGTCCCAGCTACTCNGGAGGCTMAGGCAGG300 G GTGGCGGGTGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGG300 G G T G GC G G G TGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGG300
 GGTGGCGGGTGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGG300 G GTGGC G G G TGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGG300 GGTGGCGGGTGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGG300 G GTGGGC G G GTGCCTGTAGTCCCAGCTACTC G G G A G G C T G A G G C A G G 300 GGTGGCGGGTGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGG300 G G T G GCGGGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGG300
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A G A A TGGGCGTGAACCC A GAATGGCGTGAACCC A GAATGGCGTGAACCC AGAATGGCGTGAACCC AGAATGGCGTGAACCC A GAATGGCGTGAACCC AGAATGGCGTGAACCC AGAATGGCGTGAACCC A GAATGGCGTGAACCC A G A A TGGC G T GAACC C AGAATGGCGTGAACCC A GAATGGCGTGAACCC A GAATGGCGTGAACCC A G A A T G G C G T GAACCC A GAATGGCGTGAACCC AGAATGGCGTGAACCC
 G G G A G G C G G A G C T T G C A G T G A G C C G A G A T C G C G 23 GGGGGCAGAGCTTGCA GTGAGCCHAGATCGTG350 G G G G G C A GAGCTTGCAGTGAGCCGAGATCGCG350 G G G G G CA A A GCTTGCAGTGAGCC GA GATCGCG350 G G G G G CA A A GCTTGCAGTGAGCCGAGATCGCG350 G G G G G C A GAGCTTGCAGTGAGCCGAGATCGCG350 GGGGGCGGAGCCTGCAGTGAGCTGAGATCGCG350 GGGGGCGGAGCCTGCAGTGAGCTGAGATCGCG350 GGGGGCGGAGCCTGCAGTGAGCTGAGATCGCG350 GGGGGCGGAGCTTGCAGTGAGCTGAGATCGCG350 GGGGGCGGAGCCTGCAGTGAGCTGAGATCGCG350 GGGGGC G GAGCCTGCAGTGAGCTGAGATCGC G 350 G G G G GC G GAGCCTGCAGTGAGCTGAGATCGCG350 G G G G GC G GAGCCTGCAGTGAGCTGAGATCGCG350 G G G G GC G GAGCCTGCAGTGAGCTGAGATCGCG350 G G G G GCGGAGCCTGCAGTGAGCCGAGATCGTG350


C CACTGCACTCCAGCCTGGGCGACAGAGCGAGACTCCGTCTCAAAAAA 287 CCACTGCACTCCAGCCTGGGCGATAG--CGAGACTCCGTCTCAAAAAAAA 398 C C A C T G C A C T C TA G C C T G G G T G A TA G CCACTGCACTCCAGCCTGGGTGATAG CCACTGCACTCCTAG C CACTGGACTCTAGCCTGGGGTGA C CACTGTACTCCAGCCTGGGCGAT CCACTGTACTCCAGCCTGGGCGAT C CACTGTACTCCAGCCTGGGCGA C CACT GCAC TCCAGCCTGGGCGA C CACTGCACTC CAGCCTGGGCGA CCACTGTACTCCAGCCTGGGCGA C CACTGCACTCCAGCCTGGGCGA C C A C T G C A C T C C A G C C T G G G C G A C C A C T G C A C T C C A G C C T G G G C G A CCACTGCACTCCAGCCTGGGCGA A G-

A G
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A G

## A G

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A G C G A G A C T C C G T C T C A A A A A A A A 398 C G A G A C T C C GTCTCAAAAAAA A 398 C GA GACTCCGTCTCAAAAAAAA 398 C G A G A C T C C G T C T C A A A A A A A A 398 C G A G C C T C C G T C T C A A A A A A A A 398 C GAGCCTCCGTCTCAAAAAAAA 398 C GAGCCTCCGTCTCAAAAAAAA 398 C G A G C C T C C G T C T C A A A A A A A A 398 C G A G C C T C C G T C T C A A A A A A A A 398 C GA G C C TCC GTCTCAAAAAAAA 398 C GAGCCTCCGTCTCAAAAAAAA 398 C G A G C C T C C G T C T C A A A A A A A A 398 C G A G C C T C C G T C T C A A A A A A A A 398 C GAGACTCCGTCTCAAAAAAAA 398

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Suzie Ref.
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KB4204
KB5404
KB5405


G G C C C T C A G G T T A T A A 444 G GCCCTCAGGTTATAA448 G G C C C T C A G G T T A TAA 442 G G C C C T C A G G T T A T A A 446 G G C C C T C A G G T T A TAA 441 G G C C C TCA G G T T A TAA 445 G G C C C T C A G G T T A T A A 445 G G C C C T C A G G T T A T A A 445 G G C C C T C A G G T T A TAA 445 G GCCCTCAGGTTATAA445 G G C C C T C A G G T T A T A A 445 G G C C C T C A G G T T A T A A 444 G GCCCTCAGGTTATAA 445 GGCCCTCAGGTTATAA445 G GCCCTCAGGTTATAA444

Figure S1. Sequence Alignment Report of Chr7 shared Alu insertion and flanking sequence. The TATA box-like sequence is conserved in all the species sequenced at this locus and is proximal to the 5 ' end of the Alu insertion (highlighted in blue). The Pol III transcription termination signal (TTTT) is also conserved in all species (highlighted in red) immediately after the 3' target site duplication (TSD). TSDs are highlighted in teal. Highlighted in pink are mutations different from the AluY subfamily consensus sequence that are shared by all investigated species and are all located in the right monomer of the element following the middle A-rich region. These coincide with diagnostic nucleotides of the AluYe5 subfamily. Orangutan-specific substitutions are highlighted in yellow. The Homininae-specific substitutions are highlighted in green. Chimpanzee / gorilla mutations are highlighted in brown and random mutations are shown in gray.

Table S1

| DNA sequences in Alignment Report |  | Common Names | Origin | ID number | 5' TATA box | 5' TSD | $\begin{array}{\|c\|} \hline \text { 3' TSD \& } \\ \text { termination signal } \\ \hline \end{array}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Species Names |  |  |  |  |  |  |
| 1 | Homo sapiens | Human | hg18 | N/A | TATAAAAA | CAGGTCTGTT | CAGGTCTGTTTT |
| 2 | Pan troglodytes | Chimpanzee | panTro2 | NS06006 | TATAAAAA | CAGGTCTGTT | CAGGTCTGTTTT |
| 3 | Pan paniscus | Bonobo | IPBIR | PR00661 | TATAAAAA | CAGGTCTGTT | CAGGTCTGTTTT |
| 4 | Pan paniscus | Bonobo | Coriell | AG05253 | TATAAAAA | CAGGTCTGTT | CAGGTCTGTTTT |
| 5 | Gorilla gorilla | Lowland gorilla | Coriell | AG05251 | TATAAAAA | CAGGTCTGTT | CAGGTCTGTTTT |
| 6 | Pongo pygmaeus abelii | Sumatran Orangutan | ponAbe2 | PR01109 ("Susie") | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 7 | Pongo pygmaeus abelii | Sumatran Orangutan | Coriell ${ }^{\text {* }}$ | PR01109 ("Susie") | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 8 | Pongo pygmaeus abelii | Sumatran orangutan | Coriell | GM06213A | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 9 | Pongo pygmaeus abelii | Sumatran orangutan | Coriell | GM04272A | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 10 | Pongo pygmaeus abelii | Sumatran orangutan | Coriell | NG12256 | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 11 | Pongo pygmaeus abelii | Sumatran orangutan | Coriell | NG06209 | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 12 | Pongo pygmaeus pygmaeus | Bornean Orangutan | Coriell | AG05252A | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 13 | Pongo pygmaeus pygmaeus | Bornean Orangutan | SDFZ | KB4204 | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 14 | Pongo pygmaeus pygmaeus | Bornean Orangutan | SDFZ | KB5404 | TATAAAAA | TAGGTCTGTT | CAGGTCTGTTTT |
| 15 | Pongo pygmaeus pygmaeus | Bornean Orangutan | SDFZ | KB5405 | TATAAAAA | CAGGTCTGTT | CAGGTCTGTTTT |

IPBIR: Integrated Primate Biomaterials and Information Resource
Coriell: Coriell Institute for Medical Research, 403 Haddon Avenue, Camden, NJ
SDFZ: San Diego Frozen Zoo,
Conservation and Research for Endangered Species (CRES)

* DNA PR01109 in conjunction with the Orangutan Genome Sequencing Project

Washington University Genome Sequencing Center, St. Louis, MO 63108

